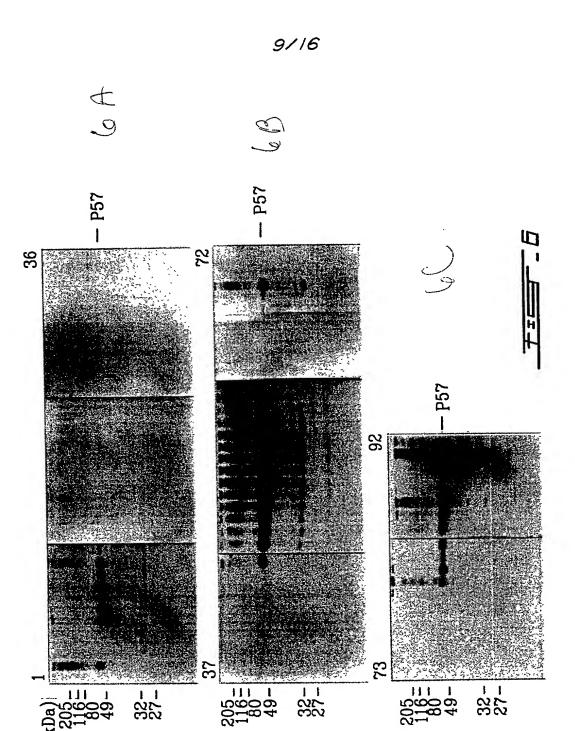
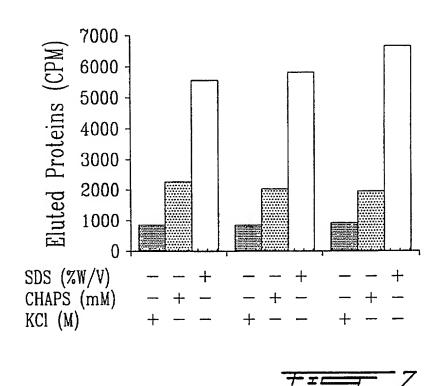
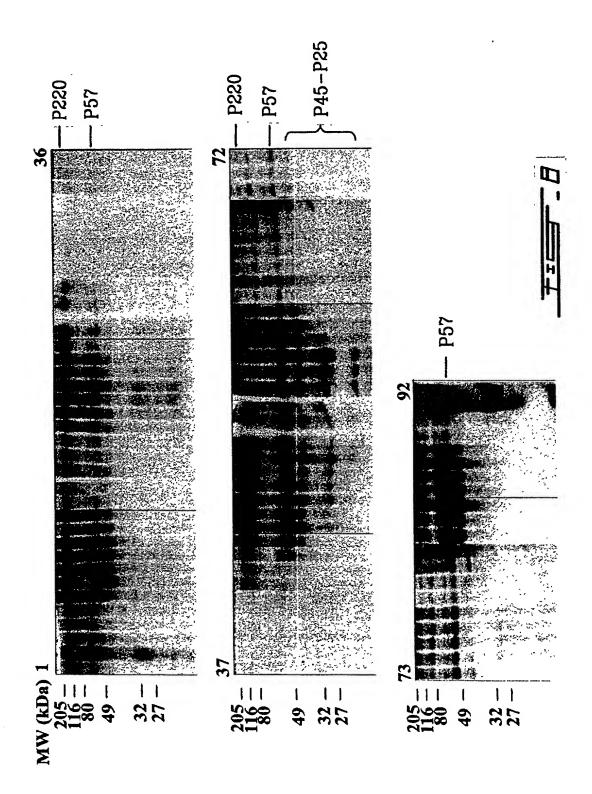


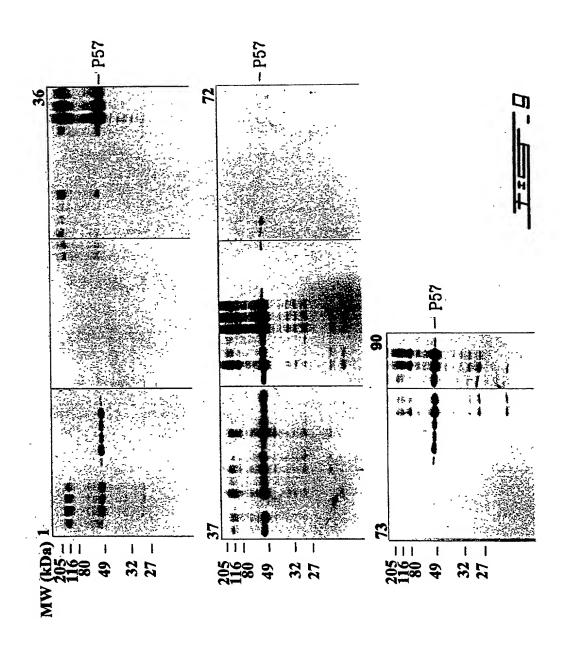
WO 00/70351 PCT/CA00/00587







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Alignment Of Amino Acid Sequences Of Human P-gp3 And P-gp1

Linker Domains

P-gp3 618 IMKKEGVYFKLVNMQTSGSQIQSEEF--ELNDEKAATRMAPNGWKSRLFR-HSTQKNLKNSQM 677

M+ N +S L R ST++++ ഥ 드 LMK++G+YFKLV MQT+G++++ E

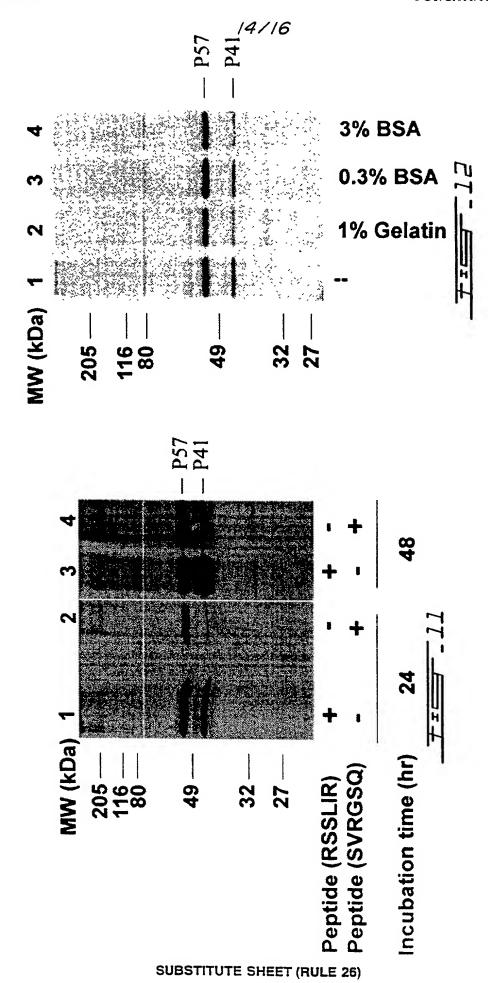
P-gp1 LMK**ekgiyfklutmo**tagnevelenaadeskseidalemssnd**srsslirkrstrrsvrgsoa** 677 615

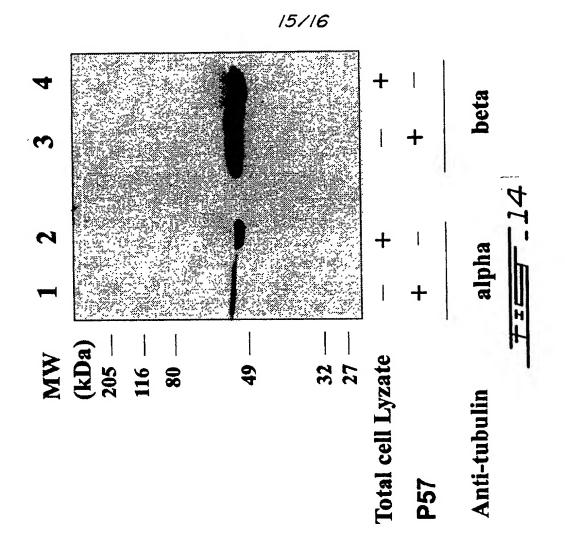
678 CQKSLDVETDGLEANVPPV**SFLKVLKLNKTEWP** 710 **P-gp3** 

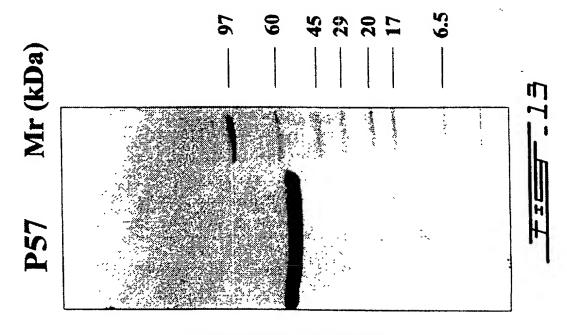
+ L + + L+ ++PPVSF +++KLN TEWP

678 QDRKLSTK-EALDESIPPV**SFWRIMKLNLTEWP** 709 P-gp1

Tz===10



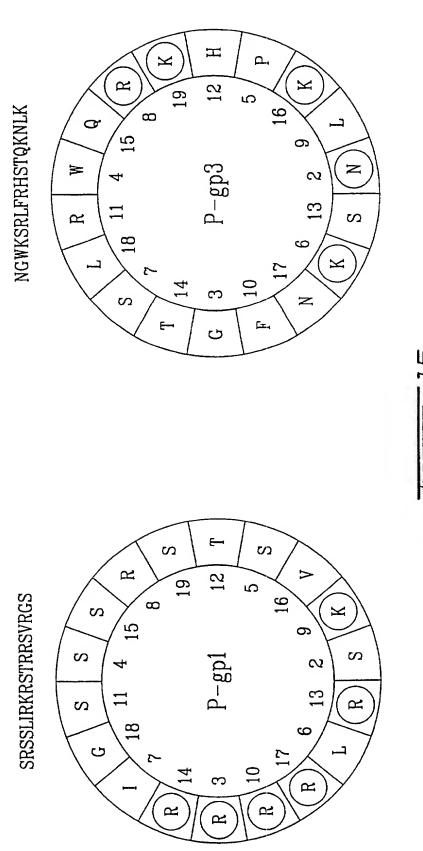




SUBSTITUTE SHEET (RULE 26)

Helical Wheel Presentation

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